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OM protein - protein search, using sw model

Run on: March 17, 2004, 18:47:23 ; Search time 25.8852 seconds
 (without alignments)
 683.183 Million cell updates/sec

Title: US-09-989-981a-4
 Perfect score: 3494

Scoring table: BIOSM62
 Sequence: 1 MAEKTKEEQLWNGVLUQDA.....FLPLVLYSLKLQKSIQDW 672

Searched: 1045404 seqs, 257433775 residues

Total number of hits satisfying chosen parameters: 1045404

Minimum DB seq length: 0
 Maximum DB seq length: 200000000

Post-processing: Minimum Match 0%
 Maximum Match 100%
 Listing first 45 summaries

Database : Published_Applications_AA:*

1: /cgn2_6/ptodata/1/pubpa/US07_PUBCOMB.pep:*

2: /cgn2_6/ptodata/1/pubpa/PCT_NEW_PUB.pep:*

3: /cgn2_6/ptodata/1/pubpa/US06_PUBCOMB.pep:*

4: /cgn2_6/ptodata/1/pubpa/US07_PUBCOMB.pep:*

5: /cgn2_6/ptodata/1/pubpa/ACTUS_OUTCOMB.pep:*

6: /cgn2_6/ptodata/1/pubpa/US08_PUBCOMB.pep:*

7: /cgn2_6/ptodata/1/pubpa/US09_PUBCOMB.pep:*

8: /cgn2_6/ptodata/1/pubpa/US08_PUBCOMB.pep:*

9: /cgn2_6/ptodata/1/pubpa/US09_PUBCOMB.pep:*

10: /cgn2_6/ptodata/1/pubpa/US09_PUBCOMB.pep:*

11: /cgn2_6/ptodata/1/pubpa/US09_PUBCOMB.pep:*

12: /cgn2_6/ptodata/1/pubpa/US09_PUBCOMB.pep:*

13: /cgn2_6/ptodata/1/pubpa/US09_PUBCOMB.pep:*

14: /cgn2_6/ptodata/1/pubpa/US09_PUBCOMB.pep:*

15: /cgn2_6/ptodata/1/pubpa/US10_PUBCOMB.pep:*

16: /cgn2_6/ptodata/1/pubpa/US10_PUBCOMB.pep:*

17: /cgn2_6/ptodata/1/pubpa/US60_NEW_PUB.pep:*

18: /cgn2_6/ptodata/1/pubpa/US60_PUBCOMB.pep:*

pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query	% Match	Length	DB	ID	Description
1	3494	100.0	672	10	US-09-989-981A-4		Sequence 4, Appli
2	2883.5	82.5	673	10	US-09-989-981A-4		Sequence 8, Appli
3	289.5	82.4	673	14	US-10-090-455-7		Sequence 7, Appli
4	1508.5	43.3	374	15	US-10-425-378-9		Sequence 9, Appli
5	753	21.6	725	12	US-10-425-599-175941		Sequence 1, Appli
6	701.5	20.1	652	9	US-09-337-927-1		Sequence 2, Appli
7	701.5	20.1	652	10	US-09-989-981A-2		Sequence 3, Appli
8	697	19.9	651	9	US-09-837-922-3		Sequence 6, Appli
9	697	19.9	651	10	US-09-989-981A-6		Sequence 14, Appli
10	697	19.9	651	14	US-10-090-455-6		Sequence 35, Appli
11	672.5	19.2	657	9	US-09-866-866A-14		Sequence 61, Appli
12	653.5	18.9	655	9	US-09-981-333-35		Sequence 1, Appli
13	659.5	18.9	655	14	US-10-405-867-61		Sequence 2, Appli
14	659.5	18.9	655	15	US-10-405-805-2		Sequence 1, Appli
15	657.5	18.8	655	10	US-09-961-086-1		Sequence 1, Appli

pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

RESULT 1
 US-09-989-981A-4
 Sequence 4, Application US/09989981A
 Publication No. US20030049730A1
 GENERAL INFORMATION:
 APPLICANT: Hobbs, Helen H.
 APPLICANT: Shan, Bei
 APPLICANT: Barnes, Robert
 APPLICANT: Tian, Hui
 APPLICANT: Tularik, Inc.
 APPLICANT: Board of Regents, The University of Texas System
 TITLE OF INVENTION: ABCG5 and ABCG8: Compositions and Methods of Use
 FILE REFERENCE: 018781-007320US
 CURRENT APPLICATION NUMBER: US 09/989, 981A
 PRIORITY APPLICATION NUMBER: US 60/252, 235
 PRIORITY FILING DATE: 2000-11-20
 PRIORITY APPLICATION NUMBER: US 60/253, 645
 PRIORITY FILING DATE: 2000-11-28
 NUMBER OF SEQ ID NOS: 13
 SOFTWARE: PatentIn Ver. 2.1
 SEQ ID NO: 4
 LENGTH: 672
 TYPE: PRT
 ORGANISM: Mus musculus
 FEATURE:
 OTHER INFORMATION: mouse ABCG8 (mABCGB)
 US-09-989-981A-4

Query Match 100.0%; Score 3494; DB 10; Length 672;
 Best Local Similarity 100.0%; Prod. No. 0; Mismatches 0; Indels 0; Caps 0;
 Matches 672; Conservative 0; Sequence 1, Appli

QY 1 MAEKTKEEQLWNGVLUQDA...GLQPSSESDNSL...YTS...GNTL...W...D...Y...Q...D...A... 60
 DQ 1 MAEKTKEEQLWNGVLUQDA...GLQPSSESDNSL...YTS...GNTL...W...D...Y...Q...D...A... 60
 61 QVFWFBQLAQKIPWRSHSQS...C...L...G...I...R...N...L...S...F...K...V...R...G...M...L...A...T...G...S...G...G...R...A...L...D...V...T...G... 120

Query Match 62.5%; Score 2883.5; DB 10; Length 673;
 Best Local Similarity 81.9%; Pred. No. 2,2e-266; Mismatches 51; Indels 1; Gaps 1;

Matches 551; Conservative 52; Mismatches 69; Indels 1; Gaps 1;

QY 121 RGHGKOMKGGQIWINQGPPSTPDLRKVYAHVHQDQLPNLTRETLAFAQMRPRTFS 180
 Db 121 RGHGKOMKGGQIWINQGPPSTPDLRKVYAHVHQDQLPNLTRETLAFAQMRPRTFS 180

QY 181 QAQDRKVEDVIALRLRQCANTRGVNTYRGVSGGERRVSVQGLNPGGDDFL 240
 Db 181 QAQDRKVEDVIALRLRQCANTRGVNTYRGVSGGERRVSVQGLNPGGDDFL 240

QY 241 SGIDSFTAINLWLTSLRAKGNLVLISLHQPSDFRLFDJULLMSGTTYLGAAQM 300
 Db 241 SGIDSFTAINLWLTSLRAKGNLVLISLHQPSDFRLFDJULLMSGTTYLGAAQM 300

QY 301 VOYFTSIGHPCPYNSPADFVDTISDRSRKEREVATEKAQSLAALFLEKQGDDFL 360
 Db 301 VOYFTSIGHPCPYNSPADFVDTISDRSRKEREVATEKAQSLAALFLEKQGDDFL 360

QY 361 WKAZAKELNTSTHTSVLTLTQDTCGAVELPGMIEOSTLRRQISNDPRLPTLHG 420
 Db 361 WKAZAKELNTSTHTSVLTLTQDTCGAVELPGMIEOSTLRRQISNDPRLPTLHG 420

QY 421 SEACIUMSLITIGFLYVGHAKQKLSMDTAALFPMGALPPNVLIDVSKCHERSMYYE 480
 Db 421 SEACIUMSLITIGFLYVGHAKQKLSMDTAALFPMGALPPNVLIDVSKCHERSMYYE 480

QY 481 LEDGLYTGPPYFAKILGSLPERCAYVITYAMPIYMLNTRPPELFLHFLWVFC 540
 Db 481 LEDGLYTGPPYFAKILGSLPERCAYVITYAMPIYMLNTRPPELFLHFLWVFC 540

QY 541 CORTMALAASAMLPATHMSSPFCNALYNSTYLTAGFMILNLTWVPAWISKSLFLNCES 600
 Db 541 CORTMALAASAMLPATHMSSPFCNALYNSTYLTAGFMILNLTWVPAWISKSLFLNCES 600

QY 601 GIMQTOFNGHLYTTOIGNFTSILGDTMISAMDINSHPELYIAVILIVISGFLFLYLIS 660
 Db 601 GIMQTOFNGHLYTTOIGNFTSILGDTMISAMDINSHPELYIAVILIVISGFLFLYLIS 660

QY 661 LKLIKQSKIONW 672
 Db 661 LKLIKQSKIONW 672

RESULT 2
 US-09-989-981A-8
 ; Sequence 8, Application US/0989981A
 ; Publication No. US20030049730A1
 ; GENERAL INFORMATION:
 ; APPLICANT: Hobbs, Helen H.
 ; APPLICANT: Shan, Bei
 ; APPLICANT: Barnes, Robert
 ; APPLICANT: Barnes, Robert
 ; APPLICANT: Tularik Inc.
 ; APPLICANT: Tularik Inc.
 ; APPLICANT: Board of Regents, The University of Texas System
 ; TITLE OF INVENTION: ABCG5 and ABCG8: Compositions and Methods of Use
 ; FILE REFERENCE: 017810-00720US
 ; CURRENT FILING DATE: 2002-07-23
 ; PRIOR APPLICATION NUMBER: US 60/252, 235
 ; PRIOR APPLICATION NUMBER: US 60/253, 645
 ; PRIOR FILING DATE: 2000-11-28
 ; NUMBER OF SEQ ID NOS: 13
 ; SOFTWARE: PatentIn Ver. 2.1
 ; SEQ ID NO 8
 ; LENGTH: 673
 ; TYPE: PRT
 ; ORGANISM: Homo sapiens
 ; FEATURE:
 ; OTHER INFORMATION: human ABCG8 (hABCG8)
 ; US-09-989-981A-8

Query Match 82.4%; Score 2879.5; DB 14; Length 673;
 ; Sequence 7, Application US/10090455
 ; Publication No. US2003027259A1
 ; GENERAL INFORMATION:
 ; APPLICANT: Chen, Hongyun
 ; APPLICANT: Le, Binh, Stephane
 ; TITLE OF INVENTION: NOVEL ABCG4 TRANSPORTER AND USES THEREOF
 ; FILE REFERENCE: 100103_406
 ; CURRENT FILING DATE: 2002-03-01
 ; NUMBER OF SEQ ID NOS: 17
 ; SOFTWARE: FASTSEQ for Windows Version 4.0
 ; SEQ ID NO 7
 ; LENGTH: 673
 ; TYPE: PRT
 ; ORGANISM: Homo sapiens
 ; US-10-090-455-7

Best Local Similarity 81.7%, Pred. No. 5.3e-266; Matches 550; Conservative 52; Mismatches 70; Indels 1; Gaps 1; Result 5

1 MARKTBETOLWNTVQDASQLODSFSSDSNLYFTSQSNLLEVDTYQDIA 60
 1 MAGKAKAERBERGPKGATPPDTSLQDRIFSSEDNSLFTYQDIA 60

61 QVPWFEOQAKKIPWRSQSSQDSCELOQIIRNISFKVRSQMLAIGGGCGRASLDVITG 120
 61 QVPWFEOQAKQFQMPWISPSQNSCELIQIUNSPKVRSQMLAIGGGCGRASLDVITG 120

121 RHHGKIKRSQGQWINGQSPSPQVLRKCVAHVROHNQIQLNLTREKQMLPFTS 180
 121 RHHGKIKRSQGQWINGQSPSPQVLRKCVAHVROHNQIQLNLTREKQMLPFTS 180

181 QAQDRKEVEDVIAELRQCANTRVQYVQGSGGRRSVIGVOLWNGLIDEP 240
 181 QAQDRKEVEDVIAELRQCANTRVQYVQGSGGRRSVIGVOLWNGLIDEP 240

181 QARQDRKEVEDVIAELRQCANTRVQYVQGSGGRRSVIGVOLWNGLIDEP 240
 181 QARQDRKEVEDVIAELRQCANTRVQYVQGSGGRRSVIGVOLWNGLIDEP 240

241 SGQDSFAHNLYTISLAKGNRLVLSHQPSDIRLFDLVLMSGTPYLGAQQM 300
 241 SGQDSFAHNLYTISLAKGNRLVLSHQPSDIRLFDLVLMSGTPYLGAQHM 300

301 VQYFTSIGHPCRYSNPADFYDVLTSIDRSKEREVATVEKAQSLAALPFLXQGFDPL 360
 301 VQYFTSIGHPCRYSNPADFYDVLTSIDRSKEREVATVEKAQSLAALPFLXQGFDPL 360

361 LWRAEAKINTSHTVSLTQD-C-GTAVELPGMIFQSTIIRQISNDFFDPL 419
 361 LWRAEAKINTSHTVSLTQD-C-GTAVELPGMIFQSTIIRQISNDFFDPL 419

361 LWRAEAKINTSHTVSLTQD-C-GTAVELPGMIFQSTIIRQISNDFFDPL 420
 361 LWRAEAKINTSHTVSLTQD-C-GTAVELPGMIFQSTIIRQISNDFFDPL 420

420 GSEACMLSLIGFLYHGAKQKLSFMDTAALLMIGALIPENVILDKVSKRSERSMLY 479
 420 GSEACMLSLIGFLYHGAKQKLSFMDTAALLMIGALIPENVILDKVSKRSERSMLY 479

421 GABACLMSTIGFLYFHGSIOLSPMDTAALLMIGALIPENVILDKVSKRSERSMLY 480
 421 GABACLMSTIGFLYFHGSIOLSPMDTAALLMIGALIPENVILDKVSKRSERSMLY 480

480 ELEDGLYTAGPFKAKIGELPEHCAVYIYAMPIYIYMLRVPFELLHELLWVYV 539
 480 ELEDGLYTAGPFKAKIGELPEHCAVYIYAMPIYIYMLRVPFELLHELLWVYV 539

481 ELEDGLYTAGPFKAKIGELPEHCAVYIYAMPIYIYMLRVPFELLHELLWVYV 540
 481 ELEDGLYTAGPFKAKIGELPEHCAVYIYAMPIYIYMLRVPFELLHELLWVYV 540

540 PCRTMALARASAMPLPTFHMSSFCNALNSFYTAGFMNLDNIVIWAWISKSFLRWC 599
 540 PCRTMALARASAMPLPTFHMSSFCNALNSFYTAGFMNLDNIVIWAWISKSFLRWC 599

541 PCRMALAAALUPTFHMSSFCNALNSFYLAGFMNLSWTVAWISKSFLRWC 600
 541 PCRMALAAALUPTFHMSSFCNALNSFYLAGFMNLSWTVAWISKSFLRWC 600

600 SGLMIOQENGHLYTQIGNFTSILGTMISAMDNLNSHPLYIYLIVGISYFLFLYU 659
 600 SGLMIOQENGHLYTQIGNFTSILGTMISAMDNLNSHPLYIYLIVGISYFLFLYU 659

601 EGLMKRIFSRRTYKMPNLNTAVSGDKLISVWELDSIPLYIYLIVGISGFMVUYV 660
 601 EGLMKRIFSRRTYKMPNLNTAVSGDKLISVWELDSIPLYIYLIVGISGFMVUYV 660

660 SLMLIKOSIQDW 672
 660 SLMLIKOSIQDW 672

661 SLMLIKOSIQDW 673
 661 SLMLIKOSIQDW 673

US-10-415-378-9

Query Match 43.2%; Score 1508.5; DB 15; Length 374; Best Local Similarity 74.9%; Pred. No. 3e-135; Mismatches 50; Indels 1; Gaps 1; Matches 260; Conservative 43; Mismatches 50; Indels 1; Gaps 1; Result 5

QY 300 VQYFTSIGHPCRYSNPADFYDVLTSIDRSKEREVATVEKAQSLAALPFLXQGFDPL 359
 QY 300 VQYFTSIGHPCRYSNPADFYDVLTSIDRSKEREVATVEKAQSLAALPFLXQGFDPL 359

Db 1 MYVFTAIGPCRYSNPADFYDVLTSIDRSKEREVATVEKAQSLAALPFLXQGFDPL 60
 Db 1 MYVFTAIGPCRYSNPADFYDVLTSIDRSKEREVATVEKAQSLAALPFLXQGFDPL 60

QY 360 LWRAEAKINTSHTVSLTQD-C-GTAVELPGMIFQSTIIRQISNDFFDPL 418
 QY 360 LWRAEAKINTSHTVSLTQD-C-GTAVELPGMIFQSTIIRQISNDFFDPL 418

Db 61 LWRAEAKINTSHTVSLTQD-C-GTAVELPGMIFQSTIIRQISNDFFDPL 120
 Db 61 LWRAEAKINTSHTVSLTQD-C-GTAVELPGMIFQSTIIRQISNDFFDPL 120

QY 419 HGBACLMSTIGFLYFHGSIOLSPMDTAALLMIGALIPENVILDKVSKRSERSMLY 478
 QY 419 HGBACLMSTIGFLYFHGSIOLSPMDTAALLMIGALIPENVILDKVSKRSERSMLY 478

Db 121 HGBACLMSTIGFLYFHGSIOLSPMDTAALLMIGALIPENVILDKVSKRSERSMLY 180
 Db 121 HGBACLMSTIGFLYFHGSIOLSPMDTAALLMIGALIPENVILDKVSKRSERSMLY 180

181 YLEDGLYTAGPFKAKIGELPEHCAVYIYAMPIYIYMLRVPFELLHELLWVYV 240
 181 YLEDGLYTAGPFKAKIGELPEHCAVYIYAMPIYIYMLRVPFELLHELLWVYV 240

QY 539 PCRTMALARASAMPLPTFHMSSFCNALNSFYTAGFMNLDNIVIWAWISKSFLRWC 598
 QY 539 PCRTMALARASAMPLPTFHMSSFCNALNSFYTAGFMNLDNIVIWAWISKSFLRWC 598

Db 241 PCRMALAAALUPTFHMSSFCNALNSFYLAGFMNLSWTVAWISKSFLRWC 300
 Db 241 PCRMALAAALUPTFHMSSFCNALNSFYLAGFMNLSWTVAWISKSFLRWC 300

QY 599 SGLMIOQENGHLYTQIGNFTSILGTMISAMDNLNSHPLYIYLIVGISYFLFLYU 658
 QY 599 SGLMIOQENGHLYTQIGNFTSILGTMISAMDNLNSHPLYIYLIVGISYFLFLYU 658

Db 301 EGLMKRIFSRRTYKMPNLNTAVSGDKLISVWELDSIPLYIYLIVGISGFMVUYV 360
 Db 301 EGLMKRIFSRRTYKMPNLNTAVSGDKLISVWELDSIPLYIYLIVGISGFMVUYV 360

659 SLMLIKOSIQDW 672
 659 SLMLIKOSIQDW 672

361 VSLRPIKQPSQDW 374
 361 VSLRPIKQPSQDW 374

US-10-424-599-175941
 ; Sequence 175941, Application US/10424599
 ; Publication No. US20040031072A1
 ; GENERAL INFORMATION:
 ; APPLICANT: La Rosa, Thomas J
 ; APPLICANT: Kovacic, David K
 ; APPLICANT: Zhou, Liwei
 ; APPLICANT: Cao, Youwei
 ; TITLE OF INVENTION: SOY Nucleic Acid Molecules and Other Molecules Associated With
 ; TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
 ; FILE REFERENCE: 38-21(53223)B

CURRENT APPLICATION NUMBER: US/10/424, 599
 CURRENT FILING DATE: 2003-04-28
 NUMBER OF SEQ ID NOS: 285684
 SEQ ID NO 175941
 LENGTH: 725; 45

TYPE: PRT
 ORGANISM: Glycine max
 FEATURE: NAME/KEY: unsure
 LOCATION: (1)..(725)
 OTHER INFORMATION: unsure at all Xaa locations

OTHER INFORMATION: Clone ID: PAT_MRT3847_129893C.1.pep
 US-10-424-599-175941

Query Match 21.6%; Score 753; DB 12; Length 725;
 Best Local Similarity 30.4%; Pred. No. 1.6e-62; Mismatches 25; Indels 42; Gaps 10;
 Matches 182; Conservative 121; N mismatches 25; N indels 42; N gaps 10;

QY 60 SQVPMPEQIAQFKPWR---SHSSDSCELGIRNLSPKRSQMLAIGSSGCRAS 113
 61 AEAFTPSKGTVPTVPTIOWRNINCSISDKSSKSRAPFLKQVNSGEAKPGRLLAIMPSGSKTT 120

QY 114 LDVITG----RGHGGKMKSGQIWINQSPSTPOLYRKCVAVHQDQLLNLTVETL 167
 121 LNLVLAGQITASPRHL--LSCYLFENGKPGSKNAYK--FAVTRQDIDFESQVLT 174

QY 168 AFTIAQMLRPTFSQDQRKVEDVIAEIRLQRQANTRYTQVGSGGERAVSIVQL 227
 175 SLATBQLPNISSAERDERFVNULFKLGLVSCADTNGDAKURGIGSGEKRLSMACEL 234

QY 228 LWNPGILIDEPTEPTSGLDSTPAHNLVTTSLRALKNRVLYLISIHPDSDIFRFDLVLMT 287
 235 LASPSVIFADEPPTGIDAFQAKYMETLQQLAQDHTVICSIHOPRGSVYKEDDILIT 294

QY 288 SGTPYIQLGAQ-QMVOYFUSINGHCPRYNPADTYDTSIDRSKEREVATVEKAOSLA 346
 295 BGSLYTAGPARDPEPLAYFSKFGYQCDPDHINPAEFLADLISIDISSADSVTSQKRPTGLV 354

QY 347 ALFLKUQGDDFLWKARAKELNTSTHTVSLTLPQDTCGATELPGM-IEPSTLRRQ 405
 355 EFSFGQR-QSAYIYATPITINDLNSRKKISOR-----AVVKKKGWWMQFXLILKRA 405

QY 406 ISNDFRDLPILHGSBACIMSLIGFLYGGAKOLSFMDTAALFFMIGALIPPNVILD 465
 406 WMOAQRDAPTPNKVRARMSIASIIFGSPVWRMGNSQTSIODRMLQVNTAMAAULTK 465

QY 466 VVSKCHERSRSRMSYLEDGTYTAKLIGSLEPEHCAVIVYIAMPWILNLRPVE 525
 466 TNGVFFKERAATDVRERAKGYYSLGIGLFLSKLAFIPIGAFFPMGAVLYPMARLHPTQ 525

QY 526 IFLHLFLVWVVFCCRTMALLAASAMLPTEHMSFFCNALYNSTYLTAGFMMINDLNIV 585
 526 RPKFGFCGIVTNEFASANGLITVGAMVPTEARAVGPSLMTVIVFGYYVNPENTPIL 585

QY 586 PAWISKLSFWRWCFSIMQFNG----HLYT"QIG----NFTPSILGTMIS 630
 586 FRWIPNVSLIWAQGLSINSBFSGLQFDHOHSFDIQTGEAQLRISFGKSRIRDIVIA 643

RESULT 6
 US-09-989-981a-4.rabp
 Sequence 1, Application US/09837992
 Patent No. US20020081657A1
 GENERAL INFORMATION:
 APPLICANT: Tian, Hui
 APPLICANT: Schultz, Joshua
 APPLICANT: Shan, Bei
 APPLICANT: Tulark Inc.
 TITLE OF INVENTION: Sterosterolemia Susceptibility Gene (SSG): Compositions
 TITLE OF INVENTION: and Methods of Use
 FILE REFERENCE: 018781-006020US
 CURRENT APPLICATION NUMBER: US/09/837,992

CURRENT FILING DATE: 2001-04-18
 PRIOR APPLICATION NUMBER: US 60/198, 465
 PRIOR FILING DATE: 2000-04-18
 NUMBER OF SEQ ID NOS: 2004, 234
 SEQ ID NO 1
 LENGTH: 655
 TYPE: PRT
 ORGANISM: Mus musculus
 FEATURE: NAME/KEY: unsure
 LOCATION: (1)..(655)
 OTHER INFORMATION: mouse sitosterolemia susceptibility gene (SSG)
 OTHER INFORMATION: amino acid sequence
 US-09-837-992-1

Query Match 20.1%; Score 701.5; DB 9; length 652;
 Best Local Similarity 29.1%; Pred. No. 1.1e-57; Mismatches 245; Indels 97; Gaps 19;
 Matches 194; Conservative 111; N mismatches 245; N indels 97; N gaps 19;

QY 24 QDSLSESSNS--LYFTYSGONTLEVRLDTYQVDAQSV-PWFEQIAQFKIPWRSHS 79
 27 QGSVAGTAREHSLSIGVLYHSV-----VSNRVGPW-----WNKIS 60

QY 80 SQDSELGIT-RNLISKVRSQSMALITGGCGGRASLIVDITGSGHGGKMKSGQIWINQ 138
 61 COQNDQDQIQLDVSLYESQIMCITGSSGKTTLDAISGRLLRTGTLGEVFNNGCE 120

QY 139 STPOLYRKCVAVHQDQLPNTRETTAFIAQMLRPTFSQDQRKVEDVIAEIRL 198
 121 LRRDQDQDCCSYVQSDVPELSLTRETRYTAMALCS-SADPYNKVEAMTELSIS 179

QY 199 OCANTRYTQVGSGGERVSIGVQIWNPGILIDEPTEPTSGLDSTPAHNLVTTSLRALKNRVLYLISIHPDSDIFRFDLVLMT 258
 200 FVADQDMIGSNGFGSSGERVSIAQQLQDPKYNMULBPTGIDCMWANQIVTLLAE 239

QY 259 AKGRNLVLTISHQPRSDIFRLFLVLTMSSTGTPITVLAQAMQVOYFTSIGHCPRYSPA 318
 240 AERDRIVIVTHQPRSELFQHDKTAKLTYGELVFCGTBEMLQPFNDGCGPCHEHSP 299

QY 319 DRYDVTISDRRSKEREVATVEKAOSLAALFLERKVQGFDLFLKZAEAKSNTSTHTVLT 378
 300 DPFYMDLTSVDTQSRRERBETKRVOMLCBKFKE----SDIYHKI-LEMIRARYKLP 353

QY 379 L---TQDTCGATELPGMIEQESTLIRQISNFRDIPTLIHLGSEACIMSLHIGF-- 432
 354 MYPFKTKDP-----PGMFGKLGVLRRVRTRNLMRNKAVIMRLVONLIMGLPFL 405

QY 433 LYGHGAKQLSPMDTAALFLMIGALIPPNVILDVSKCISERSMSYLEDGTYTAKLIGS 525
 406 LRVQNTLKGAVDORYGLKQVLAGATPTYGMVNAVNLFPMLRASDQESODGLYHKWQML 465

QY 493 FAKLGLBLPEHCAVIVYIAMPWILNLRPVEF---LL-HFLFLWLVVFCCRTM 544
 466 LAVLHLFLPSVATIPTVSSCYWTLGIVPEVARFGYPSAALLAFLPHLGBFL----TL 519

QY 545 ATAAASAMLPTEHMSFFCNALYNSTYLTAGFMMINDLNIVPAWISKLSFWRWCFSIMQFNG----HLYT"QIG----NFTPSILGTMIS 604
 520 VULGIVQNPNTI-VNSIVALLSISLGSILGSGPFRNQOEMPPKLIKGYFTFQKCYCILWV 578

QY 605 IOPNGLHYLTYQ-GNFTPSILGTMISAMDLNISHPLVAYIYLIVIGSY----- 651
 579 NEFYGL-----NFTCGGSNTSML----NHPWCA---ITQGVQFIEKTCPGATSRFT 622

QY 652 -GLFLY 657
 DB 623 ANFLILY 629

; GENERAL INFORMATION:
; APPLICANT: Robbs, Helen H.
; APPLICANT: Shan, Bei
; APPLICANT: Barnes, Robert
; APPLICANT: Tuan, Hui
; APPLICANT: Board of Regents, The University of Texas System
; TITLE OF INVENTION: ABCG5 and ABCG8: Compositions and Methods of Use
; FILE REFERENCE: 018781-007320US
; CURRENT APPLICATION NUMBER: US/09/989, 981A
; CURRENT FILING DATE: 2002-07-23
; PRIORITY APPLICATION NUMBER: US 60/252, 235
; PRIORITY FILING DATE: 2000-11-20
; NUMBER OF SEQ ID NOS: 13
; SEQ ID NO 2
; LENGTH: 652
; TYPE: PRT
; ORGANISM: Mus musculus
; FEATURE:
; OTHER INFORMATION: mouse ABCG5 (mABCG5)
; US-09-989-981A-2

Query Match 20.1%; Score 701.5; DB 10; Length 652;
Best Local Similarity 29.1%; Pred. No. 1.1e-57;
Matches 194; Conservative 131; Mismatches 245; Indels 97; Gaps 19;

Qy 24 QDSLSSRSNS--LYFTYQGSNTLEVRLDLYQDIAQV-PWFEOLAQQPKLIPWRSHS 79
Db 27 QSVTGTEEARHSLGVLHVSYS-----VSNGPWN-----WNIKS 60

Qy 80 SODSCSLGI-RNLUSFKYRGOMALIGGSSGRASLIDVTRGRHGGKMSGOIWINQDOP 138
Db 61 COOKWDQILKQVSLVIESGQIMCIGGGKTTDAGSRURRTGTLGEVFVNCE 120

Qy 139 SPDPOLVVKCVAHVRQHQDQQLNUTVETLAFIAQMLPRPSQOQRDKRVEVDIAELLR 198
Db 121 LRQDQFQCFSTVQLOSVFVLSSLTIVETRLRTYAMLAICRS-SADFYNKKVVEWTELS 179

Qy 199 QCANTRGNTTYRGSGGERRSVQVQWNGPGLIDEPSTSGLDSFRAHNVTTSLR 258
Db 180 HYADOMIGSYNFGGSSGERURVSIQALQDQPKVMDPFTGQDMCTANQVULLABL 239

Qy 259 AKGNRULVLSHQPRSDIFRFLDPUVLMSTCPIVAGAQMQVQYFTSGHPRPRYNSA 318
Db 240 ARDRIVITVTHOPRSLEFHDKLTYCBLVFCGTPBMLGPFFNNGCPCBHSNP 299

Qy 319 DFDVUDLSIDRSKRSKEREVATVKAQSLAALLEKQGDDFLWKABAKSLNTSTHTVLT 378
Db 300 DRYNDLSSVDPQSRERBETYKRVOMLECAKE----SDVYHK-LENTERARYKTLP 353

Qy 379 L---TQTDGTAPELPGMMEQFSTIIRROISNDFRDLPFTLHNGSEACMSLIIIG-- 432
Db 354 MPPFKTKDP-----PGMCKGLVIRRTRNLRNQKAVMLVQMLMGLFLITYL 405

Qy 433 LYVGHAKQSLSMDTAALIFMIGALLPPNVLVDVSKHRSRSMVYELDGLUTAGYPF 492
Db 406 LRVQNTLTKGAVTQDVGULYQDQGAPYTMGMLAVNLPMRANSQESQDGLHVKWQML 465

Qy 493 FAXTIGELPEHCAVYTYAMPYIWLNLRPVPEL-----LL--HFLLWLVVFCCRM 514
Db 456 LAVLHVLPPSVATIVFSSVQYKINGLYPVARYCQYFSALLPHLIGE-----TL 519

Qy 312 PRYSNPAFDYVLTSDRSKRSKEREVATVKAQSLAALLEKQGDDFLWKABAKLNTS 371
Db 292 PEHSNPFDFMDLSDVTDOSKRSKERIETSGRVQMSVESAVKSA----ICHKLNIRM 345

Qy 372 THTVSITL-----TQTDGTAPELPGMMEQFSTIIRROISNDFRDLPFTLHNGSEACMS 427
Db 346 KHLKLPWNPFFKTKOS-----PGVFSKLGVLRRVTRNLVANKLAVTRLQNLIM 397

Qy 428 LITGFLYXGRGAKQ-----SEMDTAALFMPGALIPPNVTDLWVKCHRSRSMLYTELDGL 485
Db 398 LFLFLFVURFTRSVNLKGAIODRVLQYQVGAATPYGMALAVNLPPVLAQSDQGGL 457

Qy 486 YTAPGPFKKAKLPEHCAVYLYAMPYIWLNLRPVPEL-----LL--HFLLWLV 537
Db 458 YQWQMLMAYALHVLPFSVWATMIFSSVQYWTGLHPEVARFGYFSAAALLPHLIGFL- 516

Query Match 19.9%; Score 697; DB 9; Length 651;
Best Local Similarity 29.1%; Pred. No. 3.1e-57;
Matches 195; Conservative 129; Mismatches 263; Indels 84; Gaps 18;

Qy 17 LODASLQDSL---FSSSDNSLYFTQGSQSNLVEFDLYQDIAQVPHFQEQLAQF 72
Db 15 LQNRGSSQSSLEGAPATAFPHSGLITLHASYSHVR-----PWNID-INSR 61

Qy 73 IFPRWSHSSQDSCSLGI-RNLUSFKYRGOMALIGGSSGRASLIDVTRGRHGGKMSQ 131
Db 62 QWTRQI-----LKDLSLVYQSGOMCINGGSSCRSTTIDAMSGRGRGATF-LGE 112

Qy 132 IWINGPSTPOLVVKCVAHVRQHQDQQLNUTVETLAFIAQMLPRPSQFQADRKVEDV 191
Db 113 VYVNGRALLRERBQFQCFSTVQLOSVFVLSSLTIVETRLRTYAMLAICRS-SADFYNKKVVEWTELS 171

Qy 192 IAEILRQCANTRGNTTYRGSGGERRSVQVQWNGPGLIDEPSTSGLDSFTAHNL 251
Db 172 MAELSHSHVADRLIGNYSLGGISOTERRVSIQALQDQPKVMDPFTGQDMCTANQI 231

Qy 252 VITLRLAKGKRVLISQPSRSDIFRFLDPUVLMSTCPIVAGAQMQVQYFTSGHPRPRYNSA 311
Db 232 VVLLVFLARARIVVLTIDOPRSelfQDkIAILSFLBLPFGTPAEMLDFFNDCCGYP 291

Qy 372 PRYSNPAFDYVLTSDRSKRSKEREVATVKAQSLAALLEKQGDDFLWKABAKLNTS 371
Db 292 PEHSNPFDFMDLSDVTDOSKRSKERIETSGRVQMSVESAVKSA----ICHKLNIRM 345

Qy 372 THTVSITL-----TQTDGTAPELPGMMEQFSTIIRROISNDFRDLPFTLHNGSEACMS 427
Db 346 KHLKLPWNPFFKTKOS-----PGVFSKLGVLRRVTRNLVANKLAVTRLQNLIM 397

Qy 428 LITGFLYXGRGAKQ-----SEMDTAALFMPGALIPPNVTDLWVKCHRSRSMLYTELDGL 485
Db 398 LFLFLFVURFTRSVNLKGAIODRVLQYQVGAATPYGMALAVNLPPVLAQSDQGGL 457

Qy 486 YTAPGPFKKAKLPEHCAVYLYAMPYIWLNLRPVPEL-----LL--HFLLWLV 537
Db 458 YQWQMLMAYALHVLPFSVWATMIFSSVQYWTGLHPEVARFGYFSAAALLPHLIGFL- 516

QY 430 IGYLYGIGAKOLSMPTPAALIFMIGALIPENVILIDVSKHS-----ERSMLYY 479
 Db 409 IGAIVFGKLNDSTGQNRKAGVLFIL-----TINQCFSSVSAVELFVVEKLFH 457
 QY 458 ELEDGLYTAGPFKATGE--IPEHCAVITYAMPIWLTNRPVBLFLFLHFLWMLV 538
 Db 518 YSASSMALAAGSOSVSVATLMLTICFVMMIFSGLVLNLTIASWL-SWQYFSIPR 575
 QY 597 WCGSGLMQIFNGHLYTQIG-----NFTSILGDTM--SAMDLNSHPIAYLIVI 647
 Db 576 YGFTALQHNEFLGONFCGLNATGNNPCNYA-TCTGEEYLVKGIDLSPWGLWKHVALA 634
 QY 648 GISYGLFLFLYLSLKLIKOS 668
 Db 635 CMIVIFLTIAVKKLFLKKY 655

RESULT 13
 US-10-120-687-61
 ; Sequence 61, Application US/10120687
 ; Publication No. US20030082155A1
 ; GENERAL INFORMATION:
 ; APPLICANT: Massachusetts General Hospital
 ; TITLE OF INVENTION: Stem Cells of the Islets of Langerhans and Their Use in Treating
 ; FILE REFERENCE: 3284/1235B
 ; CURRENT FILING DATE: US/10/120, 687
 ; PRIOR APPLICATION NUMBER: US/0/165082
 ; PRIOR FILING DATE: 1993-2-06
 ; PRIOR APPLICATION NUMBER: US 09/963, 875
 ; PRIOR FILING DATE: 2001-09-25
 ; PRIOR APPLICATION NUMBER: US 60/215109
 ; PRIOR FILING DATE: 2000-06-28
 ; PRIOR APPLICATION NUMBER: US 60/238880
 ; PRIOR FILING DATE: 2000-10-06
 ; PRIOR APPLICATION NUMBER: US 09/731261
 ; PRIOR FILING DATE: 2000-12-06
 ; SOFTWARE: Patentin version 3.1
 ; SEQ ID NO: 61
 ; LENGTH: 655
 ; TYPE: PRT
 ; ORGANISM: Homo sapiens
 ; US-10-120-687-61

Query Match 18.9%; Score 659.5; DB 14; Length 655;
 Best Local Similarity 27.2%; Pred. No. 1.2e-53; Mismatches 185; Conservative 141; Indels 85; Gaps 21;
 Matches 185; ;

QY 28 FSSESDNSL-YFTYSGSNTLEVRDLTYQDIASQVWPEQLAQKIPWRSHSQDSCEL 86
 Db 20 FPATASNDLKRAFT--EGAVIFSPHNCYRVLKSGF-----LPCRKPEVKI-- 63
 QY 87 GIRNLSFKVSGQMLAIGSGCGGRASLUDVITGKGKMKSGQIWINQSPSTPQLVRK 146
 Db 64 -LSNINGIMKPG-LNAILGTGGKSSLVLAARKDPGSL-SDGVLNGAPRPNF--K 118
 QY 147 C-VAHVRQHDLPLNUTRETLAFTAQMRLPRITSQAQRDKRVEDVIAEGLRQCANTRV 205
 Db 119 CNSGKIVNQDDWVGMGIVTVERNLQFSALRLATWMMHNERINRIVQDGLDKRDSKY 178
 QY 206 GNTYYFVGSGERRVSYGQVLLWPGILIDEPPTSGSLSTAHNTTSLRAGNRY 265
 Db 179 GTOFRGFGVSGGERKRSIGMHLITPSIILDEPTGGLDSSTANVLLKRSKGRTI 238

RESULT 14
 US-10-405-806-2
 ; Sequence 2, Application US/10405806
 ; Publication No. US20030232362A1
 ; GENERAL INFORMATION:
 ; APPLICANT: KOMATANI, HIDEYA
 ; APPLICANT: HARA, YOSHIAKU
 ; APPLICANT: KOTANI, HIDEHITO
 ; APPLICANT: NAKAGAWA, RINAKO
 ; TITLE OF INVENTION: DRUG RESISTANT GENE AND USE THEREOF
 ; FILE REFERENCE: 23495US0CONT
 ; CURRENT APPLICATION NUMBER: US/10/405, 806
 ; CURRENT FILING DATE: 2003-04-03
 ; PRIOR APPLICATION NUMBER: PCT/JP01/08112
 ; PRIOR FILING DATE: 2001-09-18
 ; PRIOR APPLICATION NUMBER: JP2000-303441
 ; PRIOR FILING DATE: 2000-10-03
 ; NUMBER OF SEQ ID NOS: 17
 ; SOFTWARE: Patentin version 3.2
 ; SEQ ID NO: 2
 ; LENGTH: 655
 ; TYPE: PRT
 ; ORGANISM: Homo sapiens
 ; US-10-405-806-2

Query Match 18.9%; Score 659.5; DB 15; Length 655;
 Best Local Similarity 27.2%; Pred. No. 1.2e-53; Mismatches 185; Conservative 141; Indels 85; Gaps 21;
 Matches 185; ;

QY 28 FSSESDNSL-YFTYSGSNTLEVRDLTYQDIASQVWPEQLAQKIPWRSHSQDSCEL 86
 Db 20 FPATASNDLKRAFT--EGAVIFSPHNCYRVLKSGF-----LPCRKPEVKI-- 63
 QY 87 GIRNLSFKVSGQMLAIGSGCGGRASLUDVITGKGKMKSGQIWINQSPSTPQLVRK 146
 Db 64 -LSNINGIMKPG-LNAILGTGGKSSLVLAARKDPGSL-SDGVLNGAPRPNF--K 118
 QY 147 C-VAHVRQHDLPLNUTRETLAFTAQMRLPRITSQAQRDKRVEDVIAEGLRQCANTRV 205
 Db 119 CNSGKIVNQDDWVGMGIVTVERNLQFSALRLATWMMHNERINRIVQDGLDKRDSKY 178
 QY 206 GNTYYFVGSGERRVSYGQVLLWPGILIDEPPTSGSLSTAHNTTSLRAGNRY 265
 Db 179 GTOFRGFGVSGGERKRSIGMHLITPSIILDEPTGGLDSSTANVLLKRSKGRTI 238

QY 266 LISMQPDRDIFRLFDLWMTSCTPINGAAGAQWQFTSIGIPCPRSNPADPYVLT 329
 Db 239 ISHQPRSIKFPLDSLTLASGRMLPHGPAEALGFFESAGYHCEAVNNPFLDII 298
 ;
 QY 326 SIDRES-----KEREVATVEK---AQSLAFLFKEVQGFDDEI-WBABAELN 369
 Db 299 NGDSTAVALNREEDKATEIEPSQDKPLIEKLAEVN---SSYKETAEHLQIS 353
 ;
 QY 370 TSHVSLSTLTDQDCTGATVPELPMIEOESTLRLRQISNDPDLPLTJHSGRACIMSLI 429
 ;
 QY 480 ELEDGLYTAGPFKATGE--IPEHCAVITYAMPIWLTNRPVBLFLFLHFLWMLV 538
 Db 518 YSASSMALAAGSOSVSVATLMLTICFVMMIFSGLVLNLTIASWL-SWQYFSIPR 575
 QY 597 WCGSGLMQIFNGHLYTQIG-----NFTSILGDTM--SAMDLNSHPIAYLIVI 647
 Db 576 YGFTALQHNEFLGONFCGLNATGNNPCNYA-TCTGEEYLVKGIDLSPWGLWKHVALA 634
 QY 648 GISYGLFLFLYLSLKLIKOS 668
 Db 635 CMIVIFLTIAVKKLFLKKY 655

RESULT 15
 US-10-120-687-61
 ; Sequence 61, Application US/10120687
 ; Publication No. US20030082155A1
 ; GENERAL INFORMATION:
 ; APPLICANT: Massachusetts General Hospital
 ; TITLE OF INVENTION: Stem Cells of the Islets of Langerhans and Their Use in Treating
 ; FILE REFERENCE: 3284/1235B
 ; CURRENT FILING DATE: 2002-04-11
 ; PRIOR APPLICATION NUMBER: US/0/165082
 ; PRIOR FILING DATE: 1993-2-06
 ; PRIOR APPLICATION NUMBER: US 09/963, 875
 ; PRIOR FILING DATE: 2001-09-25
 ; PRIOR APPLICATION NUMBER: US 60/215109
 ; PRIOR FILING DATE: 2000-06-28
 ; PRIOR APPLICATION NUMBER: US 60/238880
 ; PRIOR FILING DATE: 2000-10-06
 ; PRIOR APPLICATION NUMBER: US 09/731261
 ; PRIOR FILING DATE: 2000-12-06
 ; SOFTWARE: Patentin version 3.1
 ; SEQ ID NO: 61
 ; LENGTH: 655
 ; TYPE: PRT
 ; ORGANISM: Homo sapiens
 ; US-10-120-687-61

Query Match 18.9%; Score 659.5; DB 14; Length 655;
 Best Local Similarity 27.2%; Pred. No. 1.2e-53; Mismatches 185; Conservative 141; Indels 85; Gaps 21;
 Matches 185; ;

QY 28 FSSESDNSL-YFTYSGSNTLEVRDLTYQDIASQVWPEQLAQKIPWRSHSQDSCEL 86
 Db 20 FPATASNDLKRAFT--EGAVIFSPHNCYRVLKSGF-----LPCRKPEVKI-- 63
 QY 87 GIRNLSFKVSGQMLAIGSGCGGRASLUDVITGKGKMKSGQIWINQSPSTPQLVRK 146
 Db 64 -LSNINGIMKPG-LNAILGTGGKSSLVLAARKDPGSL-SDGVLNGAPRPNF--K 118
 QY 147 C-VAHVRQHDLPLNUTRETLAFTAQMRLPRITSQAQRDKRVEDVIAEGLRQCANTRV 205
 Db 119 CNSGKIVNQDDWVGMGIVTVERNLQFSALRLATWMMHNERINRIVQDGLDKRDSKY 178
 QY 206 GNTYYFVGSGERRVSYGQVLLWPGILIDEPPTSGSLSTAHNTTSLRAGNRY 265
 Db 179 GTOFRGFGVSGGERKRSIGMHLITPSIILDEPTGGLDSSTANVLLKRSKGRTI 238

QY 430 IGYLYGIGAKOLSMPTPAALIFMIGALIPENVILIDVSKHS-----ERSMLYY 479
 Db 409 IGAIVFGKLNDSTGQNRKAGVLFIL-----TINQCFSSVSAVELFVVEKLFH 457
 QY 458 ELEDGLYTAGPFKATGE--IPEHCAVITYAMPIWLTNRPVBLFLFLHFLWMLV 538
 Db 518 YSASSMALAAGSOSVSVATLMLTICFVMMIFSGLVLNLTIASWL-SWQYFSIPR 575
 QY 597 WCGSGLMQIFNGHLYTQIG-----NFTSILGDTM--SAMDLNSHPIAYLIVI 647
 Db 576 YGFTALQHNEFLGONFCGLNATGNNPCNYA-TCTGEEYLVKGIDLSPWGLWKHVALA 634
 QY 648 GISYGLFLFLYLSLKLIKOS 668
 Db 635 CMIVIFLTIAVKKLFLKKY 655

RESULT 16
 US-10-405-806-2
 ; Sequence 2, Application US/10405806
 ; Publication No. US20030232362A1
 ; GENERAL INFORMATION:
 ; APPLICANT: KOMATANI, HIDEYA
 ; APPLICANT: HARA, YOSHIAKU
 ; APPLICANT: KOTANI, HIDEHITO
 ; APPLICANT: NAKAGAWA, RINAKO
 ; TITLE OF INVENTION: DRUG RESISTANT GENE AND USE THEREOF
 ; FILE REFERENCE: 23495US0CONT
 ; CURRENT APPLICATION NUMBER: US/10/405, 806
 ; CURRENT FILING DATE: 2003-04-03
 ; PRIOR APPLICATION NUMBER: PCT/JP01/08112
 ; PRIOR FILING DATE: 2001-09-18
 ; PRIOR APPLICATION NUMBER: JP2000-303441
 ; PRIOR FILING DATE: 2000-10-03
 ; NUMBER OF SEQ ID NOS: 17
 ; SOFTWARE: Patentin version 3.2
 ; SEQ ID NO: 2
 ; LENGTH: 655
 ; TYPE: PRT
 ; ORGANISM: Homo sapiens
 ; US-10-405-806-2

Query Match 18.9%; Score 659.5; DB 15; Length 655;
 Best Local Similarity 27.2%; Pred. No. 1.2e-53; Mismatches 185; Conservative 141; Indels 85; Gaps 21;
 Matches 185; ;

QY 28 FSSESDNSL-YFTYSGSNTLEVRDLTYQDIASQVWPEQLAQKIPWRSHSQDSCEL 86
 Db 20 FPATASNDLKRAFT--EGAVIFSPHNCYRVLKSGF-----LPCRKPEVKI-- 63
 QY 87 GIRNLSFKVSGQMLAIGSGCGGRASLUDVITGKGKMKSGQIWINQSPSTPQLVRK 146
 Db 64 -LSNINGIMKPG-LNAILGTGGKSSLVLAARKDPGSL-SDGVLNGAPRPNF--K 118
 QY 147 C-VAHVRQHDLPLNUTRETLAFTAQMRLPRITSQAQRDKRVEDVIAEGLRQCANTRV 205
 Db 119 CNSGKIVNQDDWVGMGIVTVERNLQFSALRLATWMMHNERINRIVQDGLDKRDSKY 178
 QY 206 GNTYYFVGSGERRVSYGQVLLWPGILIDEPPTSGSLSTAHNTTSLRAGNRY 265
 Db 179 GTOFRGFGVSGGERKRSIGMHLITPSIILDEPTGGLDSSTANVLLKRSKGRTI 238

QY 266 LISMQPDRDIFRLFDLWMTSCTPINGAAGAQWQFTSIGIPCPRSNPADPYVLT 329
 Db 239 ISHQPRSIKFPLDSLTLASGRMLPHGPAEALGFFESAGYHCEAVNNPFLDII 298
 ;
 QY 326 SIDRES-----KEREVATVEK---AQSLAFLFKEVQGFDDEI-WBABAELN 369
 Db 299 NGDSTAVALNREEDKATEIEPSQDKPLIEKLAEVN---SSYKETAEHLQIS 353
 ;
 QY 370 TSHVSLSTLTDQDCTGATVPELPMIEOESTLRLRQISNDPDLPLTJHSGRACIMSLI 429
 ;
 QY 480 ELEDGLYTAGPFKATGE--IPEHCAVITYAMPIWLTNRPVBLFLFLHFLWMLV 538
 Db 518 YSASSMALAAGSOSVSVATLMLTICFVMMIFSGLVLNLTIASWL-SWQYFSIPR 575
 QY 597 WCGSGLMQIFNGHLYTQIG-----NFTSILGDTM--SAMDLNSHPIAYLIVI 647
 Db 576 YGFTALQHNEFLGONFCGLNATGNNPCNYA-TCTGEEYLVKGIDLSPWGLWKHVALA 634
 QY 648 GISYGLFLFLYLSLKLIKOS 668
 Db 635 CMIVIFLTIAVKKLFLKKY 655

RESULT 17
 US-10-405-806-2
 ; Sequence 2, Application US/10405806
 ; Publication No. US20030232362A1
 ; GENERAL INFORMATION:
 ; APPLICANT: KOMATANI, HIDEYA
 ; APPLICANT: HARA, YOSHIAKU
 ; APPLICANT: KOTANI, HIDEHITO
 ; APPLICANT: NAKAGAWA, RINAKO
 ; TITLE OF INVENTION: DRUG RESISTANT GENE AND USE THEREOF
 ; FILE REFERENCE: 23495US0CONT
 ; CURRENT APPLICATION NUMBER: US/10/405, 806
 ; CURRENT FILING DATE: 2003-04-03
 ; PRIOR APPLICATION NUMBER: PCT/JP01/08112
 ; PRIOR FILING DATE: 2001-09-18
 ; PRIOR APPLICATION NUMBER: JP2000-303441
 ; PRIOR FILING DATE: 2000-10-03
 ; NUMBER OF SEQ ID NOS: 17
 ; SOFTWARE: Patentin version 3.2
 ; SEQ ID NO: 2
 ; LENGTH: 655
 ; TYPE: PRT
 ; ORGANISM: Homo sapiens
 ; US-10-405-806-2

Query Match 18.9%; Score 659.5; DB 15; Length 655;
 Best Local Similarity 27.2%; Pred. No. 1.2e-53; Mismatches 185; Conservative 141; Indels 85; Gaps 21;
 Matches 185; ;

QY 28 FSSESDNSL-YFTYSGSNTLEVRDLTYQDIASQVWPEQLAQKIPWRSHSQDSCEL 86
 Db 20 FPATASNDLKRAFT--EGAVIFSPHNCYRVLKSGF-----LPCRKPEVKI-- 63
 QY 87 GIRNLSFKVSGQMLAIGSGCGGRASLUDVITGKGKMKSGQIWINQSPSTPQLVRK 146
 Db 64 -LSNINGIMKPG-LNAILGTGGKSSLVLAARKDPGSL-SDGVLNGAPRPNF--K 118
 QY 147 C-VAHVRQHDLPLNUTRETLAFTAQMRLPRITSQAQRDKRVEDVIAEGLRQCANTRV 205
 Db 119 CNSGKIVNQDDWVGMGIVTVERNLQFSALRLATWMMHNERINRIVQDGLDKRDSKY 178
 QY 206 GNTYYFVGSGERRVSYGQVLLWPGILIDEPPTSGSLSTAHNTTSLRAGNRY 265
 Db 179 GTOFRGFGVSGGERKRSIGMHLITPSIILDEPTGGLDSSTANVLLKRSKGRTI 238

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Db 119 CNSGGVVQDDVWGLTNTRENQFSAAIRLATTTMHEKNERINRVYQELGIDKADSKV 178
QY 206 GNTYTRGVSGGERRRVSVIGVOLLNPGILIDEPTEGSDSTFAHNUTTSRLAKGNRLV 265
Db 64 -LSNIGMKPG-INAIGPTGQKSSIDVLAARKOPSGL-SGDVINGAARRPANF-K 118
QY 179 GTOFRGVSGERKRTS1GMLITDPS1FLD3PTGDSSTANAVULLKRMSKORTI 238
Db 147 C-VAVVRQHDLQLENLTRETLAIAQRLPRTFSQARDKRVEDVIAEIRLRCANTRV 205
QY 119 CNSGVVQDDVWGLTNTRENQFSAAIRLATTTMHEKNERINRVYQELGIDKADSKV 178
Db 266 LISLHQPSDIFRLFDLVLMSGTP1VLAQAOQMVOFTS1GHP1PPRISNPADFYVLT 325
QY 239 IFSHOPRISIFKFLDS1LASSRLMHPGAQEAQYFEEAGYHCEAYNNPARDFLDII 298
Db 326 SIDRS-----KEREVATEK---AOSLAALFLEKVQGFDPL-WKAEMEKLN 369
QY 299 NGSTAVALNREEDPKATB1IEPSKQDPLIEKLAETVN---SSFYKETKAELHOLS 353
Db 370 TSHWVSLTITQDTCGAVELPGMIEQSTLRRQISNDRDPLTTLING3BACMSLI 429
QY 354 GGEKKKKITVFKESYTS---FCHQLRWVSKRSFKNLGNPQASIAQ1IVVVLGV 408
Db 430 IGFLYVGHGAKQLSFMDTAAFLMIGALIPPNVFLDVSKCHS-----ERSMYY 479
QY 409 IGAIVFGKLNDSTG1QNRAVGLFL-----TNNQCSVSSAVELFVWEKKPHTH 457
Db 480 BLEDGLYTAGPYFAKIGB-LPBCAVV1YAMPIWLTNPVPEFLFLHFLWLVV 538
QY 458 BYISGYYRVSYFLGKSLDILPMLPSI1FTCIVVFLMLGLRKADAFFVMMTLMVA 517
QY 539 FCCRTMALARASAMPTFHMSSFCNALNYSVFLTAFGMINLW-WVPAWISKLFLR 596
Db 518 YSASSMALARAAQSVVSVATLMTICVFMMIFSGLVLNLTITASWL-SW1QYFSIPR 575
QY 597 WCFSGLMQ1QFNQHLYTTOIG-----NFTES1GQDWTI-SAMDANSHPLA1YLV 647
Db 576 YGFTALQHREFLQGONFCPGLNATGNNPNCVYA-TCTGEBYLWKGOD1SPWGLWKNHVALA 634
QY 648 GISYGFLEFLYLSLKLKOKS 668
Db 635 CMIVIFLTIAYKLFLKKS 655

RESULT 15
US-09-961-086-1

; Sequence 1, Application US/0961086
; Publication No. US20030036645A1
; GENERAL INFORMATION:
; APPLICANT: UNIVERSITY OF MARYLAND, BALTIMORE
; APPLICANT: ROSS, Douglas D.
; APPLICANT: DOYLE, L. Austin
; APPLICANT: ABRUZZO, Lynne
; TITLE OF INVENTION: BREAST CANCER RESISTANCE PROTEIN (BCRP) AND THE DNA
; CURRENT APPLICATION NUMBER: US/09/961,086
; FILE REFERENCE: EP1376-019
; CURRENT FILING DATE: 2001-09-21
; PRIOR APPLICATION NUMBER: US 60/073,763
; PRIOR FILING DATE: 1998-02-05
; PRIOR FILING DATE: 1999-02-05
; NUMBER OF SEQ ID NOS: 7
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 1
; LENGTH: 655
; TYPE: PRT
; ORGANISM: Homo sapiens

Db 87 GIRLUSFKURSGOMLAITGGSGGRASLUDVITGRGHGKMGQKQWINGQESTPQVTK 146
QY 179 GTOFRGVSGERKRTS1GMLITDPS1FLD3PTGDSSTANAVULLKRMSKORTI 238
Db 147 C-VAVVRQHDLQLENLTRETLAIAQRLPRTFSQARDKRVEDVIAEIRLRCANTRV 205
QY 119 CNSGVVQDDVWGLTNTRENQFSAAIRLATTTMHEKNERINRVYQELGIDKADSKV 178
Db 266 LISLHQPSDIFRLFDLVLMSGTP1VLAQAOQMVOFTS1GHP1PPRISNPADFYVLT 325
QY 239 IFSHOPRISIFKFLDS1LASSRLMHPGAQEAQYFEEAGYHCEAYNNPARDFLDII 298
Db 326 SIDRS-----KEREVATEK---AOSLAALFLEKVQGFDPL-WKAEMEKLN 369
QY 299 NGSTAVALNREEDPKATB1IEPSKQDPLIEKLAETVN---SSFYKETKAELHOLS 353
Db 370 TSHWVSLTITQDTCGAVELPGMIEQSTLRRQISNDRDPLTTLING3BACMSLI 429
QY 354 GGEKKKKITVFKESYTS---FCHQLRWVSKRSFKNLGNPQASIAQ1IVVVLGV 408
Db 430 IGFLYVGHGAKQLSFMDTAAFLMIGALIPPNVFLDVSKCHS-----ERSMYY 479
QY 409 IGAIVFGKLNDSTG1QNRAVGLFL-----TNNQCSVSSAVELFVWEKKPHTH 457
Db 480 BLEDGLYTAGPYFAKIGB-LPBCAVV1YAMPIWLTNPVPEFLFLHFLWLVV 538
QY 458 BYISGYYRVSYFLGKSLDILPMLPSI1FTCIVVFLMLGLRKADAFFVMMTLMVA 517
QY 539 FCCRTMALARASAMPTFHMSSFCNALNYSVFLTAFGMINLW-WVPAWISKLFLR 596
Db 518 YSASSMALARAAQSVVSVATLMTICVFMMIFSGLVLNLTITASWL-SW1QYFSIPR 575
QY 597 WCFSGLMQ1QFNQHLYTTOIG-----NFTES1GQDWTI-SAMDANSHPLA1YLV 647
Db 576 YGFTALQHREFLQGONFCPGLNATGNNPNCVYA-TCTGEBYLWKGOD1SPWGLWKNHVALA 634
QY 648 GISYGFLEFLYLSLKLKOKS 668
Db 635 CMIVIFLTIAYKLFLKKS 655

Search completed: March 17, 2004, 19:53:50
Job time : 27.8852 secs

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